AB\_AYE Code

nohup IntaRNA -t AB\_AYE\_1.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_1\_targets.txt &> nohuphsa-AB\_AYE\_1.out &

nohup IntaRNA -t AB\_AYE\_2.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_2\_targets.txt &> nohuphsa-AB\_AYE\_2.out &

nohup IntaRNA -t AB\_AYE\_3.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_3\_targets.txt &> nohuphsa-AB\_AYE\_3.out &

nohup IntaRNA -t AB\_AYE\_4.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_4\_targets.txt &> nohuphsa-AB\_AYE\_4.out &

nohup IntaRNA -t AB\_AYE\_5.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_5\_targets.txt &> nohuphsa-AB\_AYE\_5.out &

nohup IntaRNA -t AB\_AYE\_6.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_6\_targets.txt &> nohuphsa-AB\_AYE\_6.out &

nohup IntaRNA -t AB\_AYE\_7.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_7\_targets.txt &> nohuphsa-AB\_AYE\_7.out &

nohup IntaRNA -t AB\_AYE\_8.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_8\_targets.txt &> nohuphsa-AB\_AYE\_8.out &

nohup IntaRNA -t AB\_AYE\_9.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_9\_targets.txt &> nohuphsa-AB\_AYE\_9.out &

nohup IntaRNA -t AB\_AYE\_10.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_10\_targets.txt &> nohuphsa-AB\_AYE\_10.out &

nohup IntaRNA -t AB\_AYE\_11.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_11\_targets.txt &> nohuphsa-AB\_AYE\_11.out &

nohup IntaRNA -t AB\_AYE\_12.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_12\_targets.txt &> nohuphsa-AB\_AYE\_12.out &

nohup IntaRNA -t AB\_AYE\_13.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_13\_targets.txt &> nohuphsa-AB\_AYE\_13.out &

nohup IntaRNA -t AB\_AYE\_14.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_14\_targets.txt &> nohuphsa-AB\_AYE\_14.out &

nohup IntaRNA -t AB\_AYE\_15.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_15\_targets.txt &> nohuphsa-AB\_AYE\_15.out &

nohup IntaRNA -t AB\_AYE\_16.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AB\_AYE\_16\_targets.txt &> nohuphsa-AB\_AYE\_16.out &